## G4G

### The National Genebank Perspective

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- genebanks for genomics
  - most presentations during this meeting: genomic analysis applied to genebank collections
    - which hypothesis?
    - what technology ?
    - what material?
    - how to manage the data?
    - what analytical tools?
    - makes sense for genebanks if the aim is building a resource (eScience)
      - technology develops too quickly
    - otherwise: genebanks should get out of it what they can
      - outputs often only show what we already know



- genomics for genebanks
  - genomics could help genebanks do their tasks more effectively
    - sample the genepools
      - determine structure of diversity
      - compare structure of diversity to structure of collection
    - conserve the material
      - duplication / redundancy
      - improvement quality passport data
      - monitoring for quality control
    - support users by optimising search strategies
      - core collections / FIGS
      - elite collections / screening pannels
      - allele mining



- what is needed to make gonomics work for genebanks?
  - genebank: questions & challenges
    - conceptual framework
      - what is it that genebanks want to improve?
    - appropriate material (SSD lines / research sets)
      - appropriately identified
  - genomics: data
    - technology appropriate to application
    - data management
  - bioinformatics & biometics: analytical tools
    - algorithms
    - visualisation



- national genebanks
  - tremendous capacity but varying quality
  - genomics generally doesn't have high priority
    - 'we're too busy keeping our material alive'
  - generally others do the genomics
    - at best genebanks collaborate with research groups
  - challenge: make genebanks benefit from genomics
    - depending on the application
    - coordination & methodology
    - capacity building (for planning and interpretation)
    - software tools (to use genomics data for genebank purposes
      -> DivSeek?)



# Thank you for your attention!



